

Figure 1

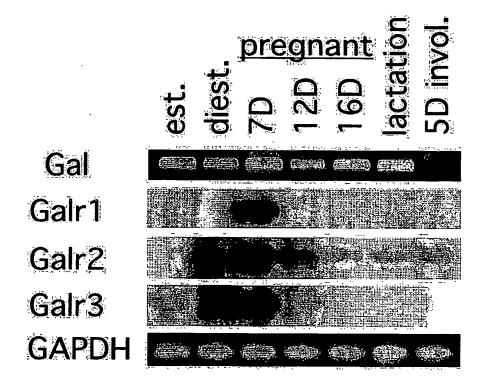


Figure 2

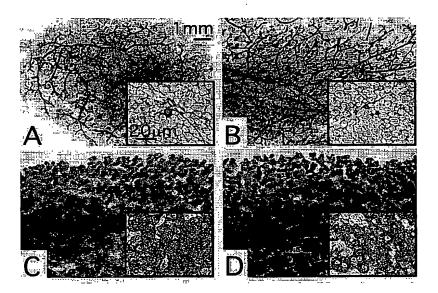


Figure 3

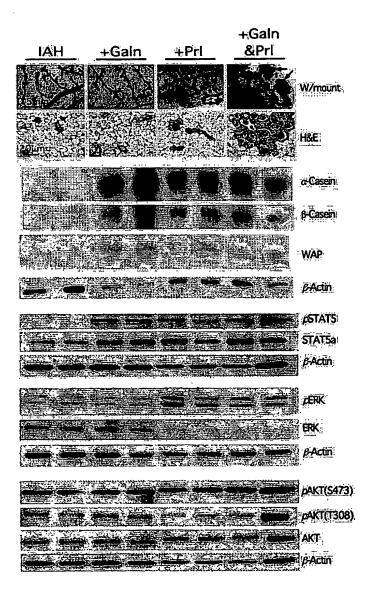


Figure 4

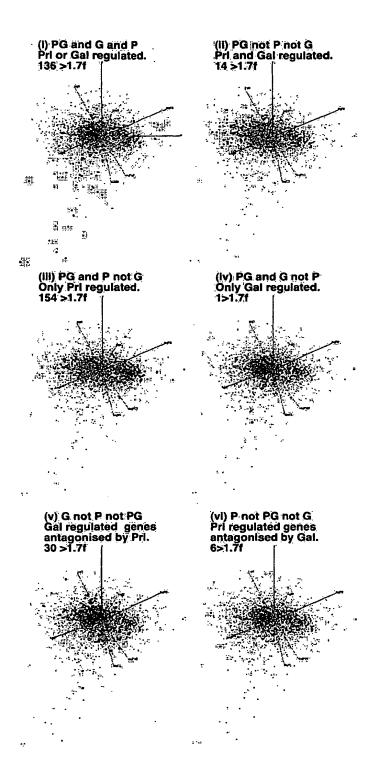


Figure 5

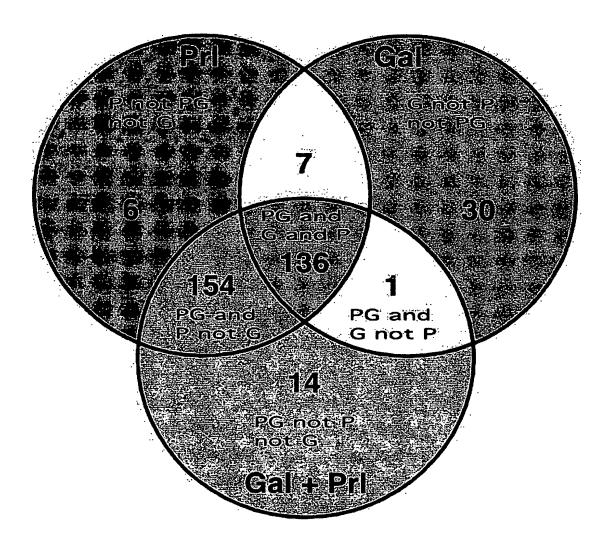


Figure 6A

				F	Fold change	ge	Ö	RT-PCR	
Accession #	Title	Symbol	Annotation	2	او	7	2	٥	- -
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V00856	idic protein		mink procein			122		2 50	257
M93428	ation depend. c	CIYCAM	cell adnesion	00	5	000			
•	Casein alpha	All Cana			2	300			
010215	casein gamma	Csng	milk protein	28	<u> </u>	8			
X04490	'casein beta	Csup	milk protein	4	64	146	106	621	200
M87863	M87863	Lactan	milk protein 🐣 🎎 🐣 🐣	.29	. 77	::31			
M10114	casein kappa	Csnk."	milk protein 📑 🕾	.26	. 33	-			
V00740	casein delta	Csnd	milk protein	414	45	42	. 19	111	129
	Cellular retinol binding protein!	CRBPI	bolism	7.4	5.7	7.1			
	E74-like factor S	+ EIFST	transcription factor	6.2	7.1	7.3	1.2	. 29	9.1
я.		WOMN		5.8	1.1	7.4	4.3	19	17
1 cil. 2 cil. 3	The second description of the second	ושטו	orowth factorial	2.3	1 6	2.7.	1.31	2.3	2.5
2007		31000	A STATE CONTROL OF THE	1,247	3 6	6 6	1.6.0.6	3.0	4.3
7.1					2 7	2.5	***		
L1244/	IGF binding protein 3	cdolb	ion signaturing		2	3	+		
M38337	globule-EGF factor 8	Mrge8	cell adhesion	2		5	٠.		
x9.2236x	M. Washing Control of the Control of	1.1SP114	Ipogenesis PH. In	12.15		5.9	7. I. Z.	3	3.4
M81445	Connexin 26	1. Cx26	gap junction	2.4	2.0	2.4			
AF077861	helix-loop-helix inhibitor, protein 1d2	142	transcription inhibitor	1.8	. 2.0	2.0.	:	·	
X16490	gen activator inhibitor 2	PAI22	protease Inhibitor,	8:51	3.5	4.4	1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1	1	
10000M	The Property of the Party of th	PAIZIA	orotease inhibitor	14.13	-1.6	-2.6			
M33300	Ben device and minimum and		t. mor-cuborascor	200	5 [6			
0.003	Cas recision gener		Tools and the second						
PG and P not o		.J			,	1			[
M19681	PDGF-Inducible protein (JE) gene	, JE	chemoattractant	4.7	• •	2:5		-	-
U03419	procollagen, type I, alpha 1	Collal	cell adhesion	2.3	7!	5.5	1	1	Ī
D29678		Coks	cell cycle			2.2			
X58251	procollagen, type I, alpha 2	Colla2	cell adhesion		4	2.2			
L42115	solute carrier family 1, member 7	Slc1a7	transport	2.0		2.0			
Y07688	nuclear factor I/X	NEX	transcription factor	1.7	1.1	1.9	-	-	
U82758	claudin 5	CldnS	tight junction	2.1	1.0	1.9			
AI154710	zinc finger protein 125	Zfp125	DNA binding	-2.2	-	-6.1	.]		
050311	myocyte enhancer factor 2B	Mef2b	transcription reg.	-2,5	-1.0	-2.0			
G not P not PG									
AV.29504400	glucose phosphate isomerase 12 complex	KKGDIJE.	CONTRACTOR SERVING TO SERVING !		7. E.	*9.1.s		4	1
AA707/8286	dermatopontin was that	CODE:	<i>7</i> 7-	*:41n3.		(F)1+3 ?	7	1	を記事が
X8115845 ASK IIGE Bindir	IGE binding to reining a set in the set in t	Figrapic .	IGE Signating Ton London	18.11.X		JE115.47	STATE OF THE PARTY		37.37.48.
WS Z683% BOLD BIRDGHS/E	eprorialphates and the sentilly		275	沙区150 6	15.230	#1F23	N. CALESCO	The state of the s	の記念
P not PG not G									
(410)0HZ	Contraction of the Contraction o	EH-BION	Cell adhesion linning to the		图6到6		Name of the last o	1	
TONS TO THE WAY TO SEE THE			protectivalenti filment		MOH W	£3285#		200	10 m
PG not P not G									
X04367	PDGF receptor beta	PDGFRB	Tyr kinase receptor	2.0	5.1-	7.5	-	-	
132838	Interleukin 1 receptor antagonist	. IL-1ra	ILR1 signaling	1.9	-2.5	2.١			
038046	topoisomerase (DNA) II beta	Top2b	DNA metabolism	1.8	1.1	-1.2	-		
X83934		Ryr3	calcium homeostasis	-2.1	-1.1	1.0	-		
AV362816	steroidogenic acute regulatory protein	Star	steroid hormone biosyn	-1.9	0.	٥.1.	-		

Figure 6B

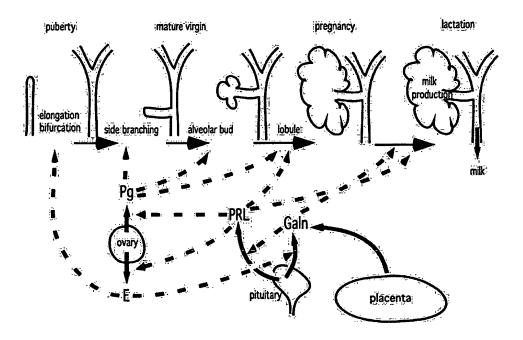


Figure 7